

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

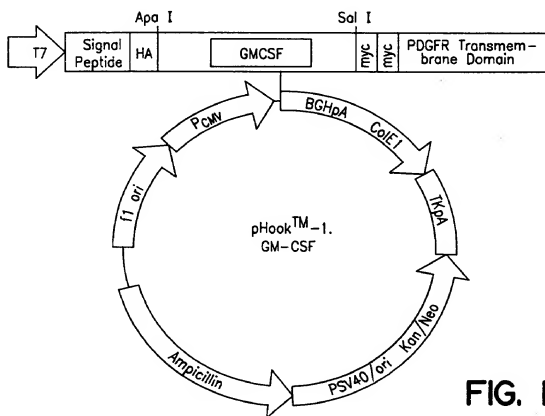


FIG. 1B

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10					20					30					40					50				
* * * *					* * * *					* * * *					* * * *					* * * *				
ATGGAGACAG					ACACACTCCT					GCTATGGGTA					CTGCTGCTCT					GGGTTCACAG				
M E T D T L L					L W V L L L T					W V P G														
<div> <div></div> <div>murine kappa light chain signal sequence</div> <div></div> </div>																								
60					70					80					90					100				
* * * *					* * * *					* * * *					* * * *					* * * *				
TTCCACTGGT					GACTATCCAT					ATGATGTTCC					AGATTATGCT					GGGGCCCAAG				
S T G D Y P					Y D V P					D Y A					G A Q									
<div> <div></div> <div></div> <div></div> </div>																								
110					120					130					140					150				
* * * *					* * * *					* * * *					* * * *					* * * *				
CACCCACCCG					CTCACCCATC					ACTGTCACCC					GGCCTTGGA					GCATGTAGAG				
A P T R S P I					T V T R P W K					H V E														
<div> <div></div> <div>mouse GMCSF</div> <div></div> </div>																								
160					170					180					190					200				
* * * *					* * * *					* * * *					* * * *					* * * *				
GCCATCAAG					AAGCCCTGAA					CCTCTGGAT					GACATGCCTG					TCACGTTGAA				
A I K E A L N					L L D D M P					V T L N														
<div> <div></div> <div></div> <div></div> </div>																								
210					220					230					240					250				
* * * *					* * * *					* * * *					* * * *					* * * *				
TGAAGAGGTA					GAAGTCGTCT					CTAACGAGTT					CTCCCTCAAG					AAGCTAACAT				
E E V E V V					S N E F					S F K					K L T									
<div> <div></div> <div></div> <div></div> </div>																								
260					270					280					290					300				
* * * *					* * * *					* * * *					* * * *					* * * *				
GTGTGCAGAC					CCGCCTGAAG					ATATTGAGC					AGGGTCTACG					GGGCAATTTC				
C V Q T R L K					I F E Q G L R					G N F														
<div> <div></div> <div></div> <div></div> </div>																								
310					320					330					340					350				
* * * *					* * * *					* * * *					* * * *					* * * *				
ACCAACTCA					AGGGCGCCTT					GAACATGACA					GCCAGCTACT					ACCAGACATA				
T K L K G A L					N M T A S Y					Y Q T Y														

FIG. 2A

APPROVED	O.G. FIG.	
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      360      370      380      390      400
      * *      * *      * *      * *      * *
CTGCCCCCA ACTCCGAAA CGGACTGTGA AACACAAGTT ACCACCTATG
  C P P    T P E    T D C E    T Q V    T T Y

      410      420      430      440      450
      * *      * *      * *      * *      * *
CGGATTTTCAT AGACAGCCTT AAAACCTTTC TGACTGATAT CCCCTTTGAA
  A D F I    D S L    K T F    L T D I    P F E

      460      470      480      490      500
      * *      * *      * *      * *      * *
TGCAAAAAAC CAGGCCAAAA AGTCGACGAA CAAAACTCA TCTCAGAAGA
  C K K    P G Q K    V D E    Q K L    I S E E
— mouse GMCSF —>

      510      520      530      540      550
      * *      * *      * *      * *      * *
GGATCTGAAT GCTGTGGGCC AGGACACGCA GGAGGTCATC GTGGTGCCAC
  D L N    A V G    Q D T Q    E V I    V V P

      560      570      580      590      600
      * *      * *      * *      * *      * *
ACTCCTTGCC CTTTAAGGTG GTGGTGATCT CAGCCATCCT GGCCCTGGTG
  H S L P    F K V    V V I    S A I L    A L V
      <— PDGFRβ transmembrane domain...—

      610      620      630      640      650
      * *      * *      * *      * *      * *
GTGCTCACCA TCATCTCCCT TATCATCCTC ATCATGCTTT GGCAGAAGAA
  V L T    I I S L    I I L    I M L    W Q K K
transmembrane domain—

      660      670      680      690
      * *      * *      * *      * *
GCCACGTTAG GCGGCCGCTC GAGATCAGCC TCGACTGTGC CTTCTAG
  P R *    A A A    R D Q P    R L C    L L
—>

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FIG. 2B

APPROVED	O.G. FIG.	
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1 2 3 4 5 6 7 8 9 10 11 12 13 14 15



FIG. 3

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
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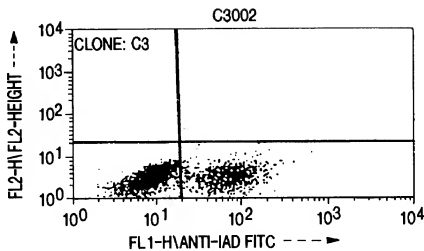


FIG. 4A

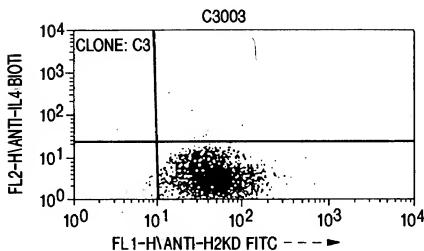


FIG. 4B

APPROVED	O.G. FIG.	
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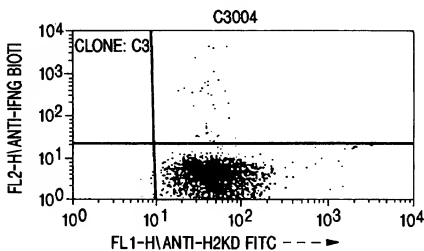


FIG. 4C

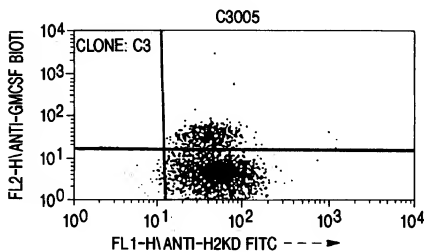


FIG. 4D